Title: COMPOSITIONS AND METHODS FOR THE THERAPY, DIAGNOSIS AND MONITORING OF BREAST CANCER Inventor(s): Gary R. Fanger et al. Serial No. 09/757,417 Docket No. 210121.479C1

WKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQVSKTEYKELLQEFIDDNATTNAIDELKECFLNQTDET**LSNVEVFNQLIYDSSLCDLF** Pro-1

MKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQVSKTEYKELLQEFIDDNATTNAI**delkecflnqtdetlsnve**vfmqliydsslcdlf **Pro-2**

MKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQVSKTEYKELLQEFIDDNATTNAIDELKECFLNQTDETLSNVEVFMQLIYDSSLCDLF Pro-3

MKLLMVLMLAALSQHCYAGSGCPLLENV**ISKTINPQVSKTEYKELLQE**FIDDNATTNAIDELKECFLNQTDETLSNVEVFMQLIYDSSLCDLF Pro-4

MKLLMVLMLAAL**SQHCYAGSGCPLLENVISKTI**NPQVSKTEYKELLQEFIDDNATTNAIDELKECFLNQTDETLSNVEVFMQLIYDSSLCDLF Pro-5

MKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQVSKTEYKELLQEFIDDNATTNAIDELKECFLNQTDETLSNVEVF**MQLIYDSSLCDLF** Pro-7

MKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQVSKTEYKELLQEFIDDNATTNAIDELKECFLNQTDET**LSNVEVFMQLIY**DSSLCDLF Pro-8

MKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQVSKTEYKELLQEFIDDNATTN**AIDELKECFL**NQTDETLSNVEVFMQLIYDSSLCDLF Pro-9

MKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQVSKTEYKELLQEFIDDNATTNAIDELKECFLNQTDETLSNVEVFMQLIYDSSLCDLF Glob-2

MKLLMVLMLAALSQHCYAGSGCPLLENVISKTINPQV**SKTEYKELLQEFIDDNATTNAIDELKECFLNQTDETLS**NVEVFMQLIYDSSLCDLF Pro-20

Mammaglobin sequence **HCYAGSGCPLLENVISK** GSGMKETAAAKFERQHMDSPDLGTDDDDKAMAISDPNS..... Peptide with Enterokinase and Thrombin cleavage sites N-terminal recombinant

Fig. 2

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	Reactivity	of Mous	Reactivity of Mouse Monoclonal		lies to Mo	ammaglobi	n with pe	ptides and	antibodies to Mammaglobin with peptides and recombinants	ıts	
Antibody	Pro2	Pro-3	Pro-4	Pro-5	Pro-6	Pro-7	Pro-8		lamma-Trx	Glob-2 Mamma-Trx N-term recomb	TRX
31-1H7	0.065	0.059	0.059	0.061	0.06	0.066	0.07	0.063	2.788	0.074	0.116
32-1611	0.026	0.022	0.054	0.054	0.055	0.057	0.055	0.055	2.75	0.057	0.07
197-1H11	0.055	0.054	0.053	1.139	0.054	0.022	0.022	0.055	2.502	2.596	0.064
304-1A5	0.054	0.054	0.053	0.053	0.054	0.053	0.023	0.054	2.7	0.056	0.064
98-1F4	0.068	0.022	0.053	0.022	0.059	0.064	0.11	0.112	2.819	0.118	0.121
296	0.055	0.057	0.056	0.056	0.055	0.62	0.026	0.637	1.566	0.069	0.159
Blank	0.056	0.055	0.053	0.055	0.052	0.053	0.053	0.053	0.056	0.052	90.0

Fig. 3A

